



PCT/09

RAW SEQUENCE LISTING

DATE: 08/26/2002

PATENT APPLICATION: US/09/831,455A

TIME: 14:40:30

Input Set : A:\pf0634usn_seqlist.txt

Output Set: N:\CRF3\08262002\I831455A.raw

2 <110> APPLICANT: INCYTE PHARMACEUTICALS, INC.
 3 TANG, Y. Tom
 4 HILLMAN, Jennifer L.
 5 YUE, Henry
 6 LAL, Preeti
 7 BANDMAN, Olga
 8 CORLEY, Neil C.
 9 GUEGLER, Karl J.
 10 BAUGHN, Mariah R.
 11 LU, Dyung Aina M.
 12 AZIMZAI, Yalda
 13 YANG, Junming
 W--> 14 <120> TITLE OF INVENTION: HUMAN HYDROLASE PROTEINS
 W--> 15 <130> FILE REFERENCE: PF-0634 PCT
 W--> 16 <140> CURRENT APPLICATION NUMBER: To Be AssignedA
 C--> 17 <141> CURRENT FILING DATE: 2002-08-26
 18 <150> PRIOR APPLICATION NUMBER: 09/190,937; unassigned; 60/135,519
 W--> 19 <151> PRIOR FILING DATE: 1998-11-12; 1998-11-12; 1999-05-21
 W--> 20 <160> NUMBER OF SEQ ID: 35
 21 <170> SOFTWARE: PERL Program
 W--> 22 <210> SEQ ID NO: 1
 23 <211> LENGTH: 159
 24 <212> TYPE: PRT
 25 <213> ORGANISM: Homo sapiens
 W--> 26 <220> FEATURE:
 27 <221> NAME/KEY: misc_feature
 28 <223> OTHER INFORMATION: Incyte ID No: 2293764CD1
 W--> 29 <400> SEQUENCE: 1
 30 Met Lys Ala Trp Gly Thr Val Val Val Thr Leu Ala Thr Leu Met
 31 1 5 10 15
 32 Val Val Thr Val Asp Ala Lys Ile Tyr Glu Leu Cys Glu Leu Ala
 33 20 25 30
 34 Ala Arg Leu Glu Arg Ala Gly Leu Asn Gly Tyr Lys Gly Tyr Gly
 35 35 40 45
 36 Val Gly Asp Trp Leu Cys Met Ala His Tyr Glu Ser Gly Phe Asp
 37 50 55 60
 38 Thr Ala Phe Val Asp His Asn Pro Asp Gly Ser Ser Glu Tyr Gly
 39 65 70 75
 40 Ile Phe Gln Leu Asn Ser Ala Trp Trp Cys Asp Asn Gly Ile Thr
 41 80 85 90
 42 Pro Thr Lys Asn Leu Cys His Met Asp Cys His Asp Leu Leu Asn
 43 95 100 105
 44 Arg His Ile Leu Asp Asp Ile Arg Cys Ala Lys Gln Ile Val Ser

Does Not Comply
Corrected Diskette Needed

pg 1, 6

Please
place
prior
application
number
directly
above
its filing
date

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45 110 115 120
 46 Ser Gln Asn Gly Leu Ser Ala Trp Thr Ser Trp Arg Leu His Cys
 47 125 130 135
 48 Ser Gly His Asp Leu Ser Glu Trp Leu Lys Gly Cys Asp Met His
 49 140 145 150
 50 Val Lys Ile Asp Pro Lys Ile His Pro
 51 155
 52 <210> SEQ ID NO: 2
 53 <211> LENGTH: 285
 54 <212> TYPE: PRT
 55 <213> ORGANISM: Homo sapiens
 W--> 56 <220> FEATURE:
 57 <221> NAME/KEY: misc_feature
 58 <223> OTHER INFORMATION: Incyte ID No: 949738CD1
 W--> 59 <400> SEQUENCE: 2
 60 Met Gly Thr Pro Gly Glu Gly Leu Gly Arg Cys Ser His Ala Leu
 61 1 5 10 15
 62 Ile Arg Gly Val Pro Glu Ser Leu Ala Ser Gly Glu Gly Ala Gly
 63 20 25 30
 64 Ala Gly Leu Pro Ala Leu Asp Leu Ala Lys Ala Gln Arg Glu His
 65 35 40 45
 66 Gly Val Leu Gly Gly Lys Leu Arg Gln Arg Leu Gly Leu Gln Leu
 67 50 55 60
 68 Leu Glu Leu Pro Pro Glu Glu Ser Leu Pro Leu Gly Pro Leu Leu
 69 65 70 75
 70 Gly Asp Thr Ala Val Ile Gln Gly Asp Thr Ala Leu Ile Thr Arg
 71 80 85 90
 72 Pro Trp Ser Pro Ala Arg Arg Pro Glu Val Asp Gly Val Arg Lys
 73 95 100 105
 74 Ala Leu Gln Asp Leu Gly Leu Arg Ile Val Glu Ile Gly Asp Glu
 75 110 115 120
 76 Asn Ala Thr Leu Asp Gly Thr Asp Val Leu Phe Thr Gly Arg Glu
 77 125 130 135
 78 Phe Phe Val Gly Leu Ser Lys Trp Thr Asn His Arg Gly Ala Glu
 79 140 145 150
 80 Ile Val Ala Asp Thr Phe Arg Asp Phe Ala Val Ser Thr Val Pro
 81 155 160 165
 82 Val Ser Gly Pro Ser His Leu Arg Gly Leu Cys Gly Met Gly Gly
 83 170 175 180
 84 Pro Arg Thr Val Val Ala Gly Ser Ser Asp Ala Ala Gln Lys Ala
 85 185 190 195
 86 Val Arg Ala Met Ala Val Leu Thr Asp His Pro Tyr Ala Ser Leu
 87 200 205 210
 88 Thr Leu Pro Asp Asp Ala Ala Ala Asp Cys Leu Phe Leu Arg Pro
 89 215 220 225
 90 Gly Leu Pro Gly Val Pro Pro Phe Leu Leu His Arg Gly Gly Gly
 91 230 235 240
 92 Asp Leu Pro Asn Ser Gln Glu Ala Leu Gln Lys Leu Ser Asp Val
 93 245 250 255

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94 Thr Leu Val Pro Val Ser Cys Ser Glu Leu Glu Lys Ala Gly Ala
 95 260 265 270
 96 Gly Leu Ser Ser Leu Cys Leu Val Leu Ser Thr Arg Pro His Ser
 97 275 280 285
 98 <210> SEQ ID NO: 3
 99 <211> LENGTH: 331
 100 <212> TYPE: PRT
 101 <213> ORGANISM: Homo sapiens
 102 <220> FEATURE:
 103 <221> NAME/KEY: misc_feature
 104 <223> OTHER INFORMATION: Incyte ID No: 1297034CD1
 105 <400> SEQUENCE: 3
 106 Met Trp Leu Trp Glu Asp Gln Gly Gly Leu Leu Gly Pro Phe Ser
 107 1 5 10 15
 108 Phe Leu Leu Leu Val Leu Leu Leu Val Thr Arg Ser Pro Val Asn
 109 20 25 30
 110 Ala Cys Leu Leu Thr Gly Ser Leu Phe Val Leu Leu Arg Val Phe
 111 35 40 45
 112 Ser Phe Glu Pro Val Pro Ser Cys Arg Ala Leu Gln Val Leu Lys
 113 50 55 60
 114 Pro Arg Asp Arg Ile Ser Ala Ile Ala His Arg Gly Gly Ser His
 115 65 70 75
 116 Asp Ala Pro Glu Asn Thr Leu Ala Ala Ile Arg Gln Ala Ala Lys
 117 80 85 90
 118 Asn Gly Ala Thr Gly Val Glu Leu Asp Ile Glu Phe Thr Ser Asp
 119 95 100 105
 120 Gly Ile Pro Val Leu Met His Asp Asn Thr Val Asp Arg Thr Thr
 121 110 115 120
 122 Asp Gly Thr Gly Arg Leu Cys Asp Leu Thr Phe Glu Gln Ile Arg
 123 125 130 135
 124 Lys Leu Asn Pro Ala Ala Asn His Arg Leu Arg Asn Asp Phe Pro
 125 140 145 150
 126 Asp Glu Lys Ile Pro Thr Leu Arg Glu Ala Val Ala Glu Cys Leu
 127 155 160 165
 128 Asn His Asn Leu Thr Ile Phe Phe Asp Val Lys Gly His Ala His
 129 170 175 180
 130 Lys Ala Thr Glu Ala Leu Lys Lys Met Tyr Met Glu Phe Pro Gln
 131 185 190 195
 132 Leu Tyr Asn Asn Ser Val Val Cys Ser Phe Leu Pro Glu Val Ile
 133 200 205 210
 134 Tyr Lys Met Arg Gln Thr Asp Arg Asp Val Ile Thr Ala Leu Thr
 135 215 220 225
 136 His Arg Pro Trp Ser Leu Ser His Thr Gly Asp Gly Lys Pro Arg
 137 230 235 240
 138 Tyr Asp Thr Phe Trp Lys His Phe Ile Phe Val Met Met Asp Ile
 139 245 250 255
 140 Leu Leu Asp Trp Ser Met His Asn Ile Leu Trp Tyr Leu Cys Gly
 141 260 265 270
 142 Ile Ser Ala Phe Leu Met Gln Lys Asp Phe Val Ser Pro Ala Tyr

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```

143          275          280          285
144 Leu Lys Lys Trp Ser Ala Lys Gly Ile Gln Val Val Gly Trp Thr
145          290          295          300
146 Val Asn Thr Phe Asp Glu Lys Ser Tyr Tyr Glu Ser His Leu Gly
147          305          310          315
148 Ser Ser Tyr Ile Thr Asp Ser Met Val Glu Asp Cys Glu Pro His
149          320          325          330

```

150 Phe

151 <210> SEQ ID NO: 4

152 <211> LENGTH: 153

153 <212> TYPE: PRT

154 <213> ORGANISM: Homo sapiens

W--> 155 <220> FEATURE:

156 <221> NAME/KEY: misc_feature

157 <223> OTHER INFORMATION: Incyte ID No: 1553276CD1

W--> 158 <400> SEQUENCE: 4

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159 Met Ala Ala Ala Leu Ala Leu Val Ala Gly Val Leu Ser Gly Ala
160 1          5          10          15
161 Val Leu Pro Leu Trp Ser Ala Leu Pro Gln Tyr Lys Lys Lys Ile
162          20          25          30
163 Thr Asp Arg Cys Phe His His Ser Glu Cys Tyr Ser Gly Cys Cys
164          35          40          45
165 Leu Met Asp Leu Asp Ser Gly Gly Ala Phe Cys Ala Pro Arg Ala
166          50          55          60
167 Arg Ile Thr Met Ile Cys Leu Pro Gln Trp Leu Glu Leu Phe Lys
168          65          70          75
169 Gly Arg Asp Cys Ile Ile Phe Ile Tyr Glu Ala Pro Thr Pro Ser
170          80          85          90
171 Leu Val Ser Ala His Asn Gln Gly Ser Tyr Gln His His Leu Pro
172          95          100          105
173 Leu Pro Asp Gly Leu Asp Val His Ile Gln Gly Leu Asp Val Phe
174          110          115          120
175 Pro Pro Val Pro Tyr Asp Leu Glu Glu Asp Ala Gly Trp Ser Leu
176          125          130          135
177 Leu Pro Trp Gly His Arg Pro Trp Leu Pro Pro Thr Cys Ser Lys
178          140          145          150

```

179 Ser Ser Ser

180 <210> SEQ ID NO: 5

181 <211> LENGTH: 571

182 <212> TYPE: PRT

183 <213> ORGANISM: Homo sapiens

W--> 184 <220> FEATURE:

185 <221> NAME/KEY: misc_feature

186 <223> OTHER INFORMATION: Incyte ID No: 1702211CD1

W--> 187 <400> SEQUENCE: 5

```

188 Met Glu Arg Ala Val Arg Val Glu Ser Gly Val Leu Val Gly Val
189 1          5          10          15
190 Val Cys Leu Leu Leu Ala Cys Pro Ala Thr Ala Thr Gly Pro Glu
191          20          25          30

```

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```

192 Val Ala Gln Pro Glu Val Asp Thr Thr Leu Gly Arg Val Arg Gly
193           35           40           45
194 Arg Gln Val Gly Val Lys Gly Thr Asp Arg Leu Val Asn Val Phe
195           50           55           60
196 Leu Gly Ile Pro Phe Ala Gln Pro Pro Leu Gly Pro Asp Arg Phe
197           65           70           75
198 Ser Ala Pro His Pro Ala Gln Pro Trp Glu Gly Val Arg Asp Ala
199           80           85           90
200 Ser Thr Ala Pro Pro Met Cys Leu Gln Asp Val Glu Ser Met Asn
201           95          100          105
202 Ser Ser Arg Phe Val Leu Asn Gly Lys Gln Gln Ile Phe Ser Val
203          110          115          120
204 Ser Glu Asp Cys Leu Val Leu Asn Val Tyr Ser Pro Ala Glu Val
205          125          130          135
206 Pro Ala Gly Ser Gly Arg Pro Val Met Val Trp Val His Gly Gly
207          140          145          150
208 Ala Leu Ile Thr Gly Ala Ala Thr Ser Tyr Asp Gly Ser Ala Leu
209          155          160          165
210 Ala Ala Tyr Gly Asp Val Val Val Val Thr Val Gln Tyr Arg Leu
211          170          175          180
212 Gly Val Leu Gly Phe Phe Ser Thr Gly Asp Glu His Ala Pro Gly
213          185          190          195
214 Asn Gln Gly Phe Leu Asp Val Val Ala Ala Leu Arg Trp Val Gln
215          200          205          210
216 Glu Asn Ile Ala Pro Phe Gly Gly Asp Leu Asn Cys Val Thr Val
217          215          220          225
218 Phe Gly Gly Ser Ala Gly Gly Ser Ile Ile Ser Gly Leu Val Leu
219          230          235          240
220 Ser Pro Val Ala Ala Gly Leu Phe His Arg Ala Ile Thr Gln Ser
221          245          250          255
222 Gly Val Ile Thr Thr Pro Gly Ile Ile Asp Ser His Pro Trp Pro
223          260          265          270
224 Leu Ala Gln Lys Ile Ala Asn Thr Leu Ala Cys Ser Ser Ser Ser
225          275          280          285
226 Pro Ala Glu Met Val Gln Cys Leu Gln Gln Lys Glu Gly Glu Glu
227          290          295          300
228 Leu Val Leu Ser Lys Lys Leu Lys Asn Thr Ile Tyr Pro Leu Thr
229          305          310          315
230 Val Asp Gly Thr Val Phe Pro Lys Ser Pro Lys Glu Leu Leu Lys
231          320          325          330
232 Glu Lys Pro Phe His Ser Val Pro Phe Leu Met Gly Val Asn Asn
233          335          340          345
234 His Glu Phe Ser Trp Leu Ile Pro Arg Gly Trp Gly Leu Leu Asp
235          350          355          360
236 Thr Met Glu Gln Met Ser Arg Glu Asp Met Leu Ala Ile Ser Thr
237          365          370          375
238 Pro Val Leu Thr Ser Leu Asp Val Pro Pro Glu Met Met Pro Thr
239          380          385          390
240 Val Ile Asp Glu Tyr Leu Gly Ser Asn Ser Asp Ala Gln Ala Lys

```

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/831,455A

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Input Set : A:\pf0634usn_seqlist.txt

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L:14 M:283 W: Missing Blank Line separator, <120> field identifier
 L:15 M:283 W: Missing Blank Line separator, <130> field identifier
 L:16 M:283 W: Missing Blank Line separator, <140> field identifier
 L:16 M:270 C: Current Application Number differs, Replaced Current Application Number
 L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date
 L:19 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
 L:20 M:283 W: Missing Blank Line separator, <160> field identifier
 L:22 M:283 W: Missing Blank Line separator, <210> field identifier
 L:26 M:283 W: Missing Blank Line separator, <220> field identifier
 L:29 M:283 W: Missing Blank Line separator, <400> field identifier
 L:56 M:283 W: Missing Blank Line separator, <220> field identifier
 L:59 M:283 W: Missing Blank Line separator, <400> field identifier
 L:102 M:283 W: Missing Blank Line separator, <220> field identifier
 L:105 M:283 W: Missing Blank Line separator, <400> field identifier
 L:155 M:283 W: Missing Blank Line separator, <220> field identifier
 L:158 M:283 W: Missing Blank Line separator, <400> field identifier
 L:184 M:283 W: Missing Blank Line separator, <220> field identifier
 L:187 M:283 W: Missing Blank Line separator, <400> field identifier
 L:269 M:283 W: Missing Blank Line separator, <220> field identifier
 L:272 M:283 W: Missing Blank Line separator, <400> field identifier
 L:324 M:283 W: Missing Blank Line separator, <220> field identifier
 L:327 M:283 W: Missing Blank Line separator, <400> field identifier
 L:358 M:283 W: Missing Blank Line separator, <220> field identifier
 L:361 M:283 W: Missing Blank Line separator, <400> field identifier
 L:415 M:283 W: Missing Blank Line separator, <220> field identifier
 L:418 M:283 W: Missing Blank Line separator, <400> field identifier
 L:465 M:283 W: Missing Blank Line separator, <220> field identifier
 L:468 M:283 W: Missing Blank Line separator, <400> field identifier
 L:538 M:283 W: Missing Blank Line separator, <220> field identifier
 L:541 M:283 W: Missing Blank Line separator, <400> field identifier
 L:566 M:283 W: Missing Blank Line separator, <220> field identifier
 L:569 M:283 W: Missing Blank Line separator, <400> field identifier
 L:598 M:283 W: Missing Blank Line separator, <220> field identifier
 L:601 M:283 W: Missing Blank Line separator, <400> field identifier
 L:656 M:283 W: Missing Blank Line separator, <220> field identifier
 L:659 M:283 W: Missing Blank Line separator, <400> field identifier
 L:750 M:283 W: Missing Blank Line separator, <220> field identifier
 L:753 M:283 W: Missing Blank Line separator, <400> field identifier
 L:800 M:283 W: Missing Blank Line separator, <220> field identifier
 L:803 M:283 W: Missing Blank Line separator, <400> field identifier
 L:868 M:283 W: Missing Blank Line separator, <220> field identifier
 L:871 M:283 W: Missing Blank Line separator, <400> field identifier
 L:889 M:283 W: Missing Blank Line separator, <220> field identifier
 L:892 M:283 W: Missing Blank Line separator, <400> field identifier
 L:918 M:283 W: Missing Blank Line separator, <220> field identifier
 L:921 M:283 W: Missing Blank Line separator, <400> field identifier
 L:962 M:283 W: Missing Blank Line separator, <220> field identifier
 L:965 M:283 W: Missing Blank Line separator, <400> field identifier

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Input Set : A:\pf0634usn_seqlist.txt

Output Set: N:\CRF3\08262002\I831455A.raw

L:979 M:283 W: Missing Blank Line separator, <220> field identifier
L:982 M:283 W: Missing Blank Line separator, <400> field identifier
L:1023 M:283 W: Missing Blank Line separator, <220> field identifier
L:1026 M:283 W: Missing Blank Line separator, <400> field identifier
L:1062 M:283 W: Missing Blank Line separator, <220> field identifier
L:1399 M:256 W: Invalid Numeric Header Field, Identifier <309> Expected, SEQ:33
L:1426 M:256 W: Invalid Numeric Header Field, Identifier <309> Expected, SEQ:34
L:1451 M:256 W: Invalid Numeric Header Field, <300> has non-blank data
L:1453 M:256 W: Invalid Numeric Header Field, Identifier <309> Expected, SEQ:35

<210> 35
 <211> 148
 <212> PRT

<213> Nasalis larvatus

<300> misc_feature

<308> GenBank ID No: g1790984

<400> 35

Met Lys Ala Leu Ile Ile Leu Gly Leu Val Leu Leu Ser Val Thr

1

5

10

15

with
response

whenever
 <308> has
 a response

(same enol
 in seqs 33-34)

It is a "header"
 only